Patent Docket P1084R1-2

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of

Paul J. Godowski et al.

Serial No.: To be assigned

Filed: 30 June 1998 (Filed Herewith)

For: ErbB4 Receptor-Specific Neuregulin

Related Ligands and Uses Therefor

Group Art Unit: Unassigned

Examiner: Unassigned

CERTIFICATION UNDER 37 CFR 1.10

EM 168 883 749 US: Express Mail Number

June 39, 1998: Date of Deposit

I hereby certify that this correspondence, consisting of CERTIFICATE RE:
SEQUENCE LISTING RESPONSE UNDER 37 CFR § 1.821(f) and (g), paper copy
and a computer-readable diskette, is being deposited with the United States
Postal Service "Express Mail Post Office to Addressee" service under 37 CFR
1.10 on the date indicated above and is addressed to the Assistant Commissioner
of Patents, Washington, D.C. 20231.

Pamela Gavette

CERTIFICATE RE: SEQUENCE LISTING

RESPONSE UNDER 37 CFR § 1.821(f) and (q)

BOX PATENT APPLICATION
Assistant Commissioner of Patents
Washington, D.C. 20231

Sir:

I hereby state that the Sequence Listing submitted herewith is submitted in paper copy and a computer-readable diskette, and that the content of the paper and computer readable copies are the same. I further state that this submission includes no new matter.

Respectfully submitted,

GENENTECH, INC.

Date: June 30, 1998

Deirdre L. Conley, Ph.D.

Reg. No. 36,487

1 DNA Way

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Phone: (650) 225-2066 Fax: (650) 952-9881 ISICYSYS LOSESSE

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RAW SEQUENCE LISTING PATENT APPLICATION US/09/107,979

DATE: 07/09/98 TIME: 13:50:58

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This Raw Listing contains the School Information Section and up to the first 5 pages.

1 SEQUENCE LISTING 2 3 (1) General Information: 4 5 (i) APPLICANT: Godowski, Paul J., Mark, Melanie Rose, Zhang, Dong Xiao 6 7 (ii) TITLE OF INVENTION: ErbB Receptor-Specific Neuregulin Related 8 Ligands and Uses Therefor 9 10 (iii) NUMBER OF SEQUENCES: 23 11 12 (iv) CORRESPONDENCE ADDRESS: 13 (A) ADDRESSEE: Genentech, Inc. 14 (B) STREET: 1 DNA Way 15 (C) CITY: South San Francisco 16 (D) STATE: California 17 (E) COUNTRY: USA 18 (F) ZIP: 94080 19 20 (v) COMPUTER READABLE FORM: 21 (A) MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk 22 (B) COMPUTER: IBM PC compatible 23 (C) OPERATING SYSTEM: PC-DOS/MS-DOS 24 (D) SOFTWARE: WinPatin (Genentech) 25 26 (vi) CURRENT APPLICATION DATA: Ø_27 (A) APPLICATION NUMBER: Unassigned (B) FILING DATE: 30-Jun-1998 28 29 (C) CLASSIFICATION: 30 31 (viii) ATTORNEY/AGENT INFORMATION: 32 (A) NAME: Conley, Deirdre L. 33 (B) REGISTRATION NUMBER: 36,487 34 (C) REFERENCE/DOCKET NUMBER: P1084R1-2 35 36 (ix) TELECOMMUNICATION INFORMATION: 37 (A) TELEPHONE: 650/225-2066 38 (B) TELEFAX: 650/952-9881 39 (2) INFORMATION FOR SEQ ID NO:1: 40 (i) SEQUENCE CHARACTERISTICS: 41 42 (A) LENGTH: 2538 base pairs 43 (B) TYPE: Nucleic Acid 44 (C) STRANDEDNESS: Single 45 (D) TOPOLOGY: Linear

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PATENT APPLICATION US/09/107,979 (ix) FEATURE: (A) NAME/KEY: mouse NRG3 nucleic acid (B) LOCATION: 1-2538 (C) IDENTIFICATION METHOD: (D) OTHER INFORMATION: (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1: CCTGACCGGC CGGCGGCCC CGGGCCGGTC TCGCCCCTCT ACCGAGCGCC 50 TCGCCGCCC CTCCCCGGCC CGCGTCCCT CCCCGTCCT CTCCTCCCCG 100 CCCGCCGCC GCCTCTCGGG GGGAGGGGCG TGGGGGCAGG GAGCCGATTT 150 GCATGCGGCC GCCGCGGCCG CTGCCTGAGC CGGAGCCCGC CGCCGCCGGA 200 GCCCGCGCCC GCGCCCGCG CCGCCCCATG CCTCTGGCGC 250 GGCCCTCGGG GGGGCGAAGG TGAAGATCGG CTCCTAGGAT GAGTGAAGGG 300 GCGGCCGGTG CCTCGCCACC TGGTGCCGCT TCGGCAGCCG CCGCCTCAGC 350

CGAGGAGGGC ACCGCGCGG CTGCGGCGGC GGCGGCGGGG GGCGGGGGGC 400
CGGACGGCGG CGGAGAAGGG GCGGCCGAAC CCCCCGGGA GTTACGCTGT 450
AGCGACTGCA TCGTGTGGAA CCGGCAGCAG ACGTGGTTGT GCGTGGTGCC 500
TCTGTTCATC GGCTTCATCG GCCTGGGGCT CAGCCTCATG CTGCTTAAAT 550
GGATCGTGGT AGGCCCGTC AAGGAGTACG TGCCCACGGA CCTGGTGGAC 600
TCCAAGGGAA TGGGCCAGGA CCCCTTCTTC CTCTCCAAGC CCAGCTCTTT 650
CCCCAAGGCT ATGGAAACCA CCACAACAAC CACTTCTACC ACGTCCCCG 700
CCACCCCCTC TGCCGGCGGC GCCGCTTCTT CCAGGACGCC TAACCGGATT 750
AGCACCCGCT TGACCACCAT CACACGGGCA CCCACCGCT TCCCTGGGCA 800
CCGGGTTCCC ATCCGGGCTA GCCCGCGCT TACCACAGCA CGGAACACTG 850
CTGCCCCTCC GACGGTCCTG TCCACCACGG CCCCTTTCTT CAGTAGCAGC 900
ACGCCCGGCT CCCGACCCCC GATGCCAGGA GCCCCCAGTA CGCAGGCGAT 950
GCCTTCCTGG CCCACTGCGG CGTATGCTAC CTCCTCTAC CTCCACGATT 1000
CCACTCCCTC CTGGACCCTG TCACCCTTC AGGATGCTGC TGCCGCCTCT 1050

TCCTCCTCAC CCTCTTCCAC CTCCTCCACT ACCACCACCC CAGAAACTAG 1100

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					L C	IF U L SL
100 101	CACCAGCCCC	AAATTTCATA	CTACAACATA	CTCCACTGAA	CGATCTGAGC	1150
102	ACTTCAAACC	CTGTCGAGAC	AAGGACCTGG	CGTATTGTCT	CAATGATGGT	1200
103 104	GAATGCTTTG	TGATTGAGAC	CCTGACAGGA	TCCCATAAGC	ACTGTCGGTG	1250
105 106	CAAGGAAGGC	TACCAAGGAG	TCCGTTGTGA	TCAATTTCTG	CCGAAAACAG	1300
107 108	АСТССАТСТТ	ATCGGATCCA	ACAGACCACT	TGGGGATTGA	ATTCATGGAG	1350
109						
110 111	AGTGAAGACG	TTTATCAAAG	GCAGGTGCTG	TCAATTTCAT	GTATCATCTT	1400
112 113	TGGAATTGTC	ATCGTGGGCA	TGTTCTGTGC	AGCATTCTAC	TTCAAAAGCA	1450
114	AGAAACAAGC	TAAACAAATT	CAGGAGCACC	TGAAAGAGTC	ACAGAATGGG	1500
115 116	AAGAACTACA	GCCTCAAGGC	ATCCAGCACA	AAGTCTGAGA	GCTTGATGAA	1550
117 118	GAGCCATGTC	CATCTACAAA	ATTATTCAAA	GGCGGATAGG	CATCCTGTGA	1600
119 120	СТССССТССА	СУУУТУТО	GAGTCAAGTT	ጥጥሮል ਫ ሮሞሮሮ	ССАСТССТТС	1650
121						
122 123	CCAGAAGTCA	CTTCTCCTGA	CCGAGGAAGC	CAGCCTATCA	AGCACCACAG	1700
124 125	CCCAGGACAA	AGGAGTGGGA	TGTTGCATAG	GAATACTTTC	AGAAGGGCAC	1750
126 127	CACCCTCACC	CCGAAGTCGA	CTGGGTGGTA	TTGTAGGACC	AGCATATCAA	1800
128	CAACTTGAAG	AATCAAGAAT	TCCAGACCAG	GATACGATAC	CTTGCCAAGG	1850
129 130	GATAGAGGTC	AGGAAGACTA	TATCCCACCT	GCCTATACAG	CTGTGGTGTG	1900
131 132	ТТСАААСАСС	ССТССАСТТА	AAGTATGTGT	ССААТСССТТ	AAGAACCCAA	1950
133						
134 135	CAAAATGCAT	CAATAAATAT	GCAACTGCCT	TCAAGAGAGA	CAAACCCCTA	2000
136 137	TTTTAATAGC	TTGGATCAAA	AGGACCTGGT	GGGTTATTTA	TCCCCAAGGG	2050
138 139	CCAATTCTGT	GCCCATCATC	CCGTCGATGG	GTCTAGAAGA	AACCTGCATG	2100
140	CAAATGCCAG	GGATTTCTGA	CGTCAAAAGC	ATTAAATGGT	GCAAAAACTC	2150
141	CTACTCCGCT	GACATTGTCA	ACGCGAGTAT	GCCAGTCAGT	GATTGTCTTC	2200
143 144	TAGAAGAACA	ACAGGAAGTG	AAAATATTAC	TAGAGACTGT	GCAGGAACAG	2250
145						
146 147	ATCCGGATTC	TGACTGATGC	CAGACGGTCA	GAAGACTTCG	AACTGGCCAG	2300
148 149	CATGGAAACT	GAGGACAGTG	CGAGCGAAAA	CACAGCCTTT	CTCCCCTGA	2350
150 151	GTCCCACGGC	CAAATCAGAA	CGAGAGGCAC	AATTTGTCTT	AAGAAATGAA	2400
152	ATACAAAGAG	ACTCTGTGCT	AACCAAGTGA	CTGGAAATGT	AGGAATCTGT	2450

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152				nu or se									
153 154	GCATTATATG CTT	IGCTAAA CAGGA	AGGAG AGGAAATTA	A ATACAAATTA 2500									
155 156	MMMamamada mma		amaam maaaaaa	0530									
157													
158	(2) INFORMATION FOR SEQ ID NO:2:												
159													
160													
161		TH: 713 amino											
162		: Amino Acid	40240										
163		LOGY: Linear											
164													
165	(ix) FEATURE:												
166	(A) NAME	KEY: Mouse N	RG3 (mNRG3)/ami	no acid seq.									
167	(B) LOCA	TION: 1-713		-									
168		TIFICATION ME											
169	(D) OTHE	R INFORMATION	:										
170													
171	(xi) SEQUENCE	DESCRIPTION:	SEQ ID NO:2:										
172													
173	_	_		o Gly Ala Ala Ser									
174	1	5	10	15									
175	310 310 310 31		Glas Glas Mbas Al										
176 177	ATA ATA ATA ATA	a Ser Ala Glu 20		a Ala Ala Ala									
178		20	25	30									
179	אות הות הות הות	. cl., cl., cl.,	Dro Aco Cl., Cl	y Gly Glu Gly Ala									
180	ATG ATG ATG ATG	35	40										
181		33	40	45									
182	Ala Glu Pro Pro	Ara Glu Leu	Ara Cue Ser Ae	p Cys Ile Val Trp									
183	014 110 110	50	55	p cys lie val lip 60									
184		•		30									
185	Asn Arg Gln Glr	Thr Trp Leu	Cvs Val Val Pr	o Leu Phe Ile Gly									
186	J	65	70	75									
187													
188	Phe Ile Gly Let	Gly Leu Ser	Leu Met Leu Le	u Lys Trp Ile Val									
189		80	85	90									
190													
191	Val Gly Ser Val	. Lys Glu Tyr	Val Pro Thr As	p Leu Val Asp Ser									
192		95	100	105									
193		_											
194	Lys Gly Met Gly			r Lys Pro Ser Ser									
195		110	115	120									
196	n)		_, _, _, _,										
197	rne Pro Lys Ala			r Thr Ser Thr Thr									
198		125	130	135									
199	Con Dro Nie Mie	Dwa C 37-	al., al., .11	- G G 1 m1									
200 201	ser Pro Ala Thi			a Ser Ser Arg Thr									
201		140	145	150									
202	Pro Asn Ara Tla	Cor The A	Iou The The Ti	e Thr Arg Ala Pro									
203	I TO MAIL MIG ITE	155	160	_									
205		133	100	165									

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206 207 208	Thr	Arg	Phe	Pro	Gly 170	His	Arg	Val	Pro	Ile 175	Arg	Ala	Ser		Arg 180
209 210 211	Ser	Thr	Thr	Ala	Arg 185	Asn	Thr	Ala	Ala	Pro 190	Pro	Thr	Val	Leu	Ser 195
212 213 214	Thr	Thr	Ala	Pro	Phe 200	Phe	Ser	Ser	Ser	Thr 205	Pro	Gly	Ser	Arg	Pro 210
215 216 217					Ala 215					220				_	225
218 219 220					Ala 230					235		_			240
221 222 223					Ser 245			•		250					255
224 225 226					Ser 260					265					270
227 228 229					Lys 275					280	-				285
230 231 232					Lys 290					295					300
233 234 235					Glu 305	_				310				_	315
236 237 238 239					Arg 320					325					330
240 241 242					Pro 335					340					345
242 243 244 245								-		355		_		_	360
246 247 248					Ser 365 Cys			_		370		_			375
249 250 251		_			380 Gln				_	385	_		_	_	390
252 253 254		7			395 Lys				-	400				_	405
255 256 257					410 His					415			٠		420
25 <i>7</i> 258	пЛр	26T	птр	νац	425	rea	GTII	ASN	ryr	430	гÀ2	АТЯ	ASP	Arg	H1S 435